

WEST Search History

DATE: Sunday, April 02, 2006

Hide?	<u>Set</u> <u>Name</u>	<u>Query</u>	<u>Hit</u> <u>Count</u>
		<i>DB=USPT; PLUR=YES; OP=OR</i>	
<input type="checkbox"/>	L33	L32	111
		<i>DB=PGPB,USPT; PLUR=YES; OP=OR</i>	
<input type="checkbox"/>	L32	GST.clm.	406
<input type="checkbox"/>	L31	GST/clm	0
<input type="checkbox"/>	L30	GST	18586
<input type="checkbox"/>	L29	RNA and L27	1
<input type="checkbox"/>	L28	Si adj RNA and L27	0
<input type="checkbox"/>	L27	6030954.pn.	1
<input type="checkbox"/>	L26	(GST OR SH1 or PTB or LIM or armadillo or Notch adj ankyrin adj repeat or zinc adj finger or leucine adj zipper or helix adj turn adj helix or helix adj loop adj helix) and 19	1
<input type="checkbox"/>	L25	(GST OR SH1 or PTB or LIM or armadillo or Notch adj ankyrin adj repeat or zinc adj finger or leucine adj zipper or helix adj turn adj helix or helix adj loop adj helix) and 11	1
<input type="checkbox"/>	L24	multivalent same recognition same GST same polypeptide	1
<input type="checkbox"/>	L23	multimer with ligand with glutathione	0
<input type="checkbox"/>	L22	multimer with ligand with GST	0
<input type="checkbox"/>	L21	(peptide or \$peptide) with (multimer or trimer or pentamer or multivalent) same helix adj turn adj helix same library	0
<input type="checkbox"/>	L20	(peptide or \$peptide) with (multimer or trimer or pentamer or multivalent) same leucine adj zipper same library	0
<input type="checkbox"/>	L19	(peptide or \$peptide) with (multimer or trimer or pentamer or multivalent) same zinc adj finger same library	0
<input type="checkbox"/>	L18	(peptide or \$peptide) with (multimer or trimer or pentamer or multivalent) same Notch adj ankyrin same library	0
<input type="checkbox"/>	L17	(peptide or \$peptide) with (multimer or trimer or pentamer or multivalent) same armadillo same library	0
<input type="checkbox"/>	L16	(peptide or \$peptide) with (multimer or trimer or pentamer or multivalent) same LIM same library	0
<input type="checkbox"/>	L15	(peptide or \$peptide) with (multimer or trimer or pentamer or multivalent) same PTB same library	0
<input type="checkbox"/>	L14	(peptide or \$peptide) with (multimer or trimer or pentamer or multivalent) same PH same library	4
<input type="checkbox"/>	L13	(peptide or \$peptide) with (multimer or trimer or pentamer or multivalent) same	0

	SH1 same library	
<input type="checkbox"/>	L12 (peptide or \$peptide) with (multimer or trimer or pentamer or multivalent) same GST same library	6
<input type="checkbox"/>	L11 (peptide or \$peptide) with (multimer or trimer or pentamer or multivalent) same GST	24
<input type="checkbox"/>	L10 (peptide or \$peptide) same (\$mer or multivalent) same GST	273
<input type="checkbox"/>	L9 6709821.pn.	1
<input type="checkbox"/>	(multivalent or multimer or biligand) same (GST OR SH1 or PTB or LIM or armadillo or Notch adj ankyrin adj repeat or zinc adj finger or leucine adj zipper or helix adj turn adj helix or helix adj loop adj helix)	848
<input type="checkbox"/>	(multivalent or multimer or biligand) same (GST OR SH1 or ph or PTB or LIM or armadillo or Notch adj ankyrin adj repeat or zinc adj finger or leucine adj zipper or helix adj turn adj helix or helix adj loop adj helix)	2386
<input type="checkbox"/>	L6 PH and L1	1
<input type="checkbox"/>	L5 SH and L1	0
<input type="checkbox"/>	L4 SH1 and L1	1
<input type="checkbox"/>	L3 GST and L1	1
<input type="checkbox"/>	L2 recognition and L1	1
<input type="checkbox"/>	L1 10/807856	1

END OF SEARCH HISTORY